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REQUEST FOR FILING APPLICATION

Under Rule 53(a), (b) & (f)

(No Filing Fee or Oath/Declaration)

(Do NOT use for Provisional or PCT Applications)

Use for Design or Utility Applications

PATENT
APPLICATION

jc772 U.S. PTO



09/20/00

RULE 53(f) NO DECLARATION

Assistant Commissioner of Patents

Atty. Dkt.

PM 273944

0421-NF

206/00

M#

Client Ref

and Trademarks
Washington, DC 20231

Date:

September 20, 2000

Sir:

1. This is a Request for filing a new Patent Application (☐ Design ☒ Utility) entitled:

2. (Complete) Title: CHIRAL, CHARGED PEPTIDE NUCLEIC ACID OLIGOMERS FROM CYCLIC MONOMERS

without a filing fee or Oath/Declaration but for which is enclosed the following:

3. ☒ Abstract 1 page(s).4. 39 Pages of Specification (only spec. and claims); 5. ☐ Specification in non-English language

6. 13 Numbered claim(s); and

7. ☒ Drawings: 13 sheet(s) ☐ 1 set informal; 8. ☒ formal of size: ☒ A4 ☐ 11"9. DOMESTIC/INTERNATIONAL priority is claimed under 35 USC 119(e)/120/365(c) based on the following provisional, nonprovisional and/or PCT international application(s):

Application No.	Filing Date	Application No.	Filing Date
(1)		(2)	
(3)		(4)	
(5)		(6)	

10. FOREIGN priority is claimed under 35 USC 119(a)-(d)/365(b) based on filing in

Application No.	Filing Date	Application No.	Filing Date
(1)		(2)	
(3)		(4)	
(5)		(6)	

11. (No.) Certified copy (copies): ☐ attached; ☐ previously filed (date) _____
in U.S. Application No. _____ / _____ filed on _____12. ☐ This is a reissue of Patent No. _____13. ☐ See top first page re prior Provisional, National, International application(s) (X box only if info is there and do not complete corresponding item 14 or 15.)14. ☐ Amend the specification by inserting before the first line -- This is a ☐ Continuation-in-Part
☐ Divisional ☐ Continuation ☐ Substitute Application (MPEP 201.09) of:14(a) ☐ National Appln. No. _____ / _____ filed .- - (M#)14(b) ☐ International Appln. No. PCT/ _____ filed _____ which designated the U.S. - -15. ☐ Amend the specification by inserting before the first line: --This application claims the benefit of U.S. Provisional Application No. 60/ ___, filed ___ --16. Extension to date: ☐ concurrently filed ☐ not needed ☐ previously filed

17. ☐ Prior application is assigned to

by Assignment recorded _____ Reel _____ Frame _____

18. ☐ Attached:

19. This application is made by the following named inventor(s) (Double check instructions for accuracy.):

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20. NOTE: FOR ADDITIONAL INVENTORS, check box ☐ and attach sheet with same information regarding additional inventors.

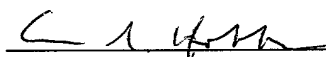
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NOTE: File in duplicate with 2 post card receipts (PAT-103) & attachments

APPLICATION UNDER UNITED STATES PATENT LAWS

Atty. Dkt. No. PM 273944

(M#)

Invention: CHIRAL, CHARGED PEPTIDE NUCLEIC ACID OLIGOMERS FROM CYCLIC MONOMERS

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000260-44795560

This is a:

- ☐ Provisional Application
- ☒ Regular Utility Application
- ☐ Continuing Application
 - ☐ The contents of the parent are incorporated by reference
- ☐ PCT National Phase Application
- ☐ Design Application
- ☐ Reissue Application
- ☐ Plant Application
- ☐ Substitute Specification
 - Sub. Spec Filed _____
 - in App. No. _____ / _____
- ☐ Marked up Specification re
 - Sub. Spec. filed _____
 - In App. No. _____ / _____

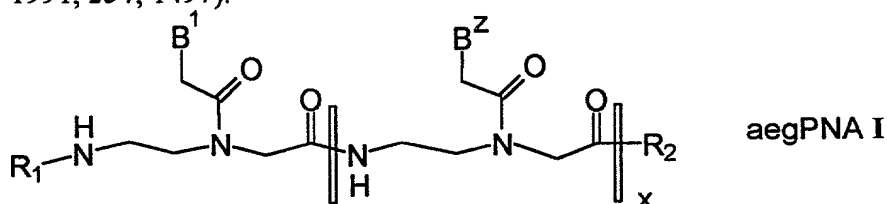
SPECIFICATION

FIELD OF THE INVENTION

This invention is related to novel compounds derived from peptide nucleic acids that bind to complementary DNA and RNA strands. In particular, the invention is concerned with peptide nucleic acid analogs that are chiral, positively charged, more soluble in aqueous systems and bind to complementary nucleic acid sequences with high avidity and sequence discrimination ability.

BACKGROUND OF THE INVENTION AND PRIOR ART

Gene targeted drugs are designed with oligonucleotides of nucleobase sequence of 10-20 units, complementary to the regulatory region of the disease associated target gene. Upon administration, they bind to specific promoter targets (DNA) and block access to RNA polymerase and as a consequence, no mRNA or the corresponding gene products are produced (Agarwal, S. in Applied antisense oligonucleotide technology, Stein, C.A. & Kreig, A.M. Ed., 1978, Wiley-Liss Inc). Depending on the target chosen, the drugs can be developed for application as antiviral, anticancer agents or other instances where conventional drugs have serious limitations. They are also useful for research and in diagnostics for detection and isolation of specific nucleic acids. Since natural oligonucleotides are degraded by nucleases, there is considerable interest in synthetic analogues of oligonucleotides that are stable under physiological conditions (Uhlmann, E. and Peyman, A. *Chem. Rev.* 1990, 90, 543; Leyten, I., Herdewijn, P., *Eur. J. Chem.*, 1998, 33, 515). In order to achieve this, a large number of variations in polynucleotide backbones have been undertaken, although so far not with the desired results. The great majority of these backbone modifications led to decreased stability for hybrids formed between the modified oligonucleotide and its complementary natural oligonucleotide, except in the case of aminoethylglycyl peptide nucleic acids (aegPNA I) (Nielsen, P. E. *et al Science* 1991, 254, 1497).



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In these compounds, a linear amide chain composed of repeating aminoethylglycine units to which nucleobase ligands are attached replaces the cyclic sugar-phosphate backbone of DNA and are amenable for easy preparation by solid phase peptide synthesis. PNAs can be used to target ss or ds DNA to produce gene regulatory molecules and as hybridization probes. Polypyrimidine PNAs form triple helices with ds DNA and are useful as antigene agents. Triplex formation and strand displacement are the only known principles in the art for sequence-specific recognition of ds DNA, with strand displacement being superior to the former. This is because unlike triplex formation, it is not restricted to homopurine-homopyrimidine sequences and allows the recognition of any sequence by Watson-Crick base pairing of the four natural nucleobases. The strand invasion of double strand DNA by unmodified PNAs is only possible in the absence of salts or at ionic strengths much lower than the physiological conditions. (Hyrup, B. and Nielsen, P.E., *BioMed. Chem. Lett.* 1996, 4, 5).

The PNA molecules although make stable hybrids with complementary DNA, are poorly soluble in aqueous media and being non-chiral bind to complementary DNA in both parallel and antiparallel directions whenever the sequence permits (Hanvey, J. C. *et al Science* 1992, 258, 1481; Uhlmann, E. *et al Angew. Chem. Int. Ed. Engl.* 1998, 37, 2796). Several structural variations within such a PNA backbone have been sought, in order to improve the water solubility, orientation specificity and DNA double strand invasion at physiological conditions, without compromising on the DNA affinity and sequence specificity. Typically, these involve conjugation with positively charged ligands such as lysine (Kim, S. H. *et al J. Am. Chem. Soc.* 1993, 115, 6477; Koch, T. *et al Tetrahedron Lett.* 1995, 36, 6933; Ishihara, T. and Corey, D. R. *J. Am. Chem. Soc.* 1999, 121, 2012) and introduction of substituents in the ethylenediamine or glycine sectors of the backbone. (Haaime, G. *et al Angew. Chem. Int. Ed. Engl.* 1996, 35, 1939; Gangamani, B. P. *et al Tetrahedron* 1999, 55, 177; D'Costa, M. *et al Organic Lett.* 1999, 1, 1513.). However, it is well understood in the art of the PNA backbone modifications that no PNA analogs have yet emerged where these problems have been simultaneously and satisfactorily addressed (Ganesh and Nielsen, *Curr. Organic Chemistry*, 2000, 9, 916). The following relevant patents on PNA so far are cited here.

	Patent No.	Authors	Title
1	WO 9964449	Harriot, P.; Nelson, J.; Wallace, A.	New cell permeable signal peptides useful for intracellular delivery of molecules
2	WO 9816550	Lowe, G.	New proline based chiral peptide nucleic acid compounds- having strong hybridisation activity, useful as antisense, antigene agents or in molecular biology
3	WO 9714793	Demidov, V. V.; Frank-Kamenetskii, M. D.; Veselkov, A. G.	Nucleic acid clamp comprising two sequences of peptide nucleic acids- connected by flexible linker with stabilizer at ends, used e.g., to cleave target nucleic acid
4	WO 9620212	Buchardt, O.; Lagriffoul, P.; Nielsen, P. E.	New peptide nucleic acid monomers with chiral backbone- may be used in preparation of oligomers useful e.g., as research tools

OBJECTS OF THE INVENTION

It is the primary object of the present invention to provide novel cyclic, chiral PNAs – homo or hetero oligomers derived from the cyclic prolyl monomers that bind ssDNA and RNA strands to form stable hybrids with much higher affinity than the parent peptide nucleic acids.

It is one object of the present invention to provide cyclic monomers for peptide nucleic acid oligomers that bind single and double strand DNAs to form stable hybrids with higher affinity, than the corresponding complexes of the standard peptide nucleic acids I composed from linear monomer units with DNA.

It is another object of invention to provide monomers that are both cyclic and chiral for making oligomers that bind one strand of a double stranded polynucleotide, in a sequence and orientation specific manner.

It is yet another object of the invention to provide cyclic, chiral monomers that confer positive charge/s on the derived oligomers and make them more soluble in water than the standard peptide nucleic acids.

It is a further object to provide synthetic procedures for cyclic, chiral peptide nucleic acid monomers needed for the oligomers of the invention.

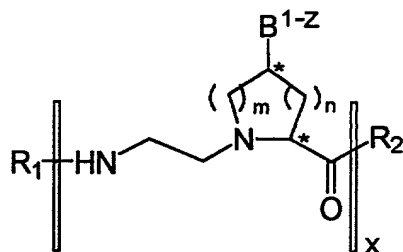
SUMMARY OF THE INVENTION

In summary, the invention provides novel entities viz., cyclic, chiral, positively charged oligomers that have specific advantages of higher solubility in water than standard PNA and bind to complementary DNA sequences in a sequence and orientation specific manner.

DETAILED DESCRIPTION OF THE INVENTION

The present invention provides a novel class of compounds (*aep*PNA II) incorporating cyclic monomers based on aminoethylprolyl (*aep*PNA) units that bind specifically to complementary ss and ds DNA and RNA strands. The compounds of the invention generally comprise ligands B (natural nucleobases thymine, cytosine, adenine or guanine, inosine, uracil and their synthetic analogs) linked to cyclic monomeric units that are an integral part of the peptide backbone. Further, the presence of two chiral carbon atoms (denoted by *) leads to four possible diastereomers and the compounds of the invention comprise all these stereomers.

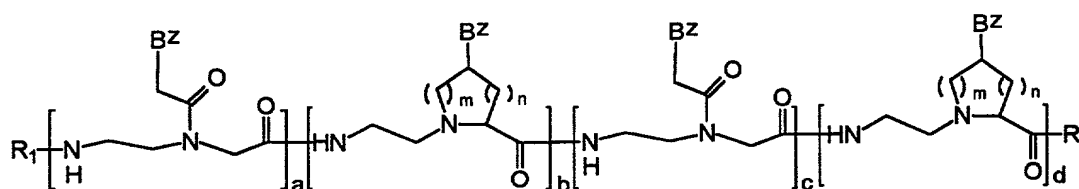
In general embodiment, the oligomers of the invention are either homopolymers *aep*PNA II of the type



aep PNA II

- Wherein m and n are 1 to 2 and x = 1-20,
- each of B¹-B^z is independently selected from the group consisting of H, HO, NH₂, naturally occurring nucleobases adenine (A), thymine (T), cytosine (C) and guanine (G), non-naturally occurring nucleobases, DNA intercalators, heterocyclic moieties and reporter ligands,
- each chiral monomeric unit independently selected from the four possible diastereomers,
- R₁=H/Fluorophore/Biotin, R₂=OH/NH(CH₂)₂COOH/
NH(CH₂)₃NH(CH₂)₄NH(CH₂)₃NH₂.

Or heteropolymers aepPNA III involving one or more substitution of the aeg monomeric unit in the aepPNA II as below :



aep PNA III

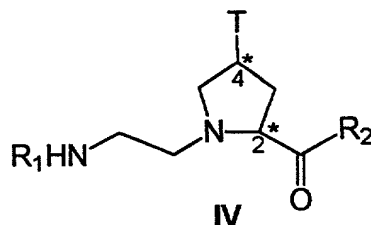
Wherein

- a, b,c,d,m,n are integers with values in the range 1to10 and various combinations thereof,
- R₁ is H/COCH₃ or L (corresponding to a fluorophore *e.g.* dansyl, carboxyfluorescein),
- R₂ is OH, NH₂, NH CH₂CH₂COOH, sperminyl. i.e., NH(CH₂)₃NH(CH₂)₄ NH(CH₂)₃NH₂.

In the following preferred embodiments, the compounds of the invention have m=n=1; B^z = T; R₁=H; R₂ = NH (CH₂)₂COOH, with

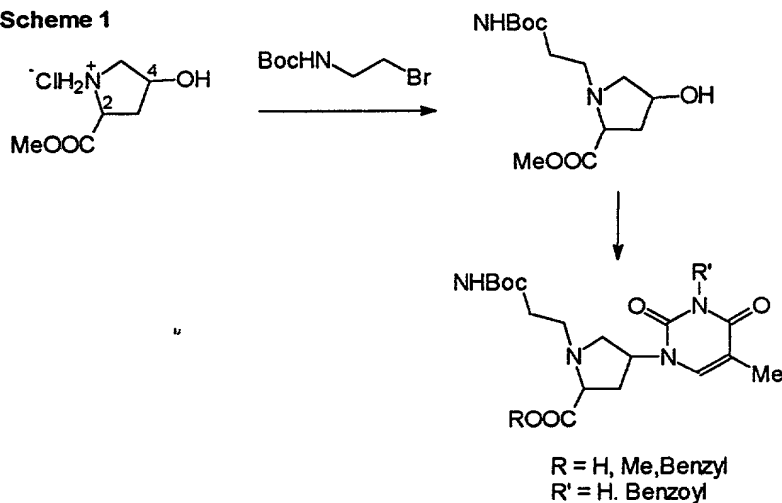
- a=7, b=1, c=d=0,
- a=c=3, b=d=1,
- a=b=c=d=1, repeating twice in that order,
- a=b=c=0, d=8, and
- a=d=0, b=1, c=7-11 and with various combinations of B^z.

The oligomers of the invention are synthesized by adaptation of standard peptide synthesis procedures, either in solution or in solid phase. The preferred monomer precursor-synthon, according to the invention, has the structure IV



Where $R_1 = \text{H/tBoc}$, $R_2 = \text{Ome/OH}$ at positions at 2 and 4 lead to four diastereomers, each synthesized by the process described in the invention (scheme 1) depending upon the stereochemistry of the starting material.

Scheme 1



In all the monomer synthons and peptide nucleic acid analogs, according to the invention, the nucleobase is T. In alternative structures, B may be a non-naturally occurring nucleobase (nucleobase analog), another base-binding moiety, an aromatic moiety, hydroxyl or even hydrogen. The homooligomers *aepPNA II*, each incorporating the above monomers at all positions or heteropolymers *aepPNA III*, with inclusion of standard peptide nucleic acid monomers in the *aepPNA II* at one or more sites, are able to recognize the complementary polypurine, single stranded DNA, generating

*aep*PNA:DNA complexes. Such a recognition of DNA taking place in the range of 8-15 bases is of biological interest.

While the oligomeric compounds of the invention show remarkably enhanced binding with complementary DNA as indicated by the increased T_m of hybrids, they also have significant discriminating ability as the presence of even a single mismatch in the complementary DNA strand would lead to significant destabilization of the complex. The present oligomeric compounds of the invention *aep*PNA II and *aep*PNA III are also more soluble in water than standard PNA I, making them more suitable for biological applications.

Whereas the improved binding of the compounds of the invention should render them efficient as antisense agents, it is expected that an extended range of related analogs or derivatives, whose structures are mentioned in the invention, may cause strand displacement. This should render them efficient antigene agents in inhibition of gene expression in cells.

BRIEF DESCRIPTION OF THE ACCOMPANYING DRAWINGS

Figure 1: Scheme for synthesis of (N-Boc)-2-aminoethylbromide

Figure 2: Scheme for N-alkylation of 4*R*-hydroxy-2*S*-prolinemethylester

Figure 3: Scheme for the synthesis of [1-(N-Boc)-2-aminoethyl]-4*S*-(N3-Benzoylthymine-1-yl)-2*S*-prolinemethylester

Figure 4: Scheme for N-alkylation of 4*R*-hydroxy-2*R*-prolinemethylester

Figure 5: Scheme for the synthesis of [1-(N-Boc)-2-aminoethyl]-4*S*-(N3-Benzoylthymine-1-yl)-2*R*-prolinemethylester

Figure 6: General scheme for ester hydrolysis to obtain protected monomer units

Figure 7: General scheme for solid phase synthesis of oligomers showing preparation of linear unprotected polyamides carrying novel monomers of the invention

Figure 8: Oligomer sequences comprising the novel monomer of the invention

Figure 9: Reverse phase HPLC of some new oligomers of the invention

Figure 10: MALDI-TOF spectra of some new oligomers of the invention

Figure 11. UV melting profiles

Figure 12: Gel electrophoresis.

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The peptide nucleic acids have been found to be superior to oligonucleotides since they have significantly higher affinity for complementary RNA or DNA. Since they are amides of non-biological amino acids, they are biostable and resistant to enzymatic degradation by proteases and nucleases. However, their properties in terms of solubility, cellular uptake and specific orientation in hybridization are not satisfactory and need to be improved by novel approaches.

The main descript of the invention is the structural design which simultaneously installs positive charges and chirality on the main backbone accompanied by controlled conformational restriction as a strategy to overcome the many limitations of peptide nucleic acids. This is achieved by linking of the α -carbon of glycyl component with the β -carbon of the side arm carrying the ligand B in peptide nucleic acids I to obtain the monomer synthons IV. The compounds of the invention, the size of the cyclic system are determined by independent values for m and n, which vary between 1 and 2. In the preferred synthons of the invention, $m=n=1$, correspond to aminoethylprolyl PNA *aep*PNA(II). An important consequence of such a structural design is the presence of a tertiary amino group in the oligomer of the invention, which has a potential for protonation at physiological pH, leading to positively charged oligomers that have better hybridization properties with the negatively charged complementary DNA and improved solubility in aqueous systems.

In another interesting aspect of the invention, chirality is introduced at 2 centers in the cyclic pyrrolidine ring system, resulting in 4 diastereomeric monomers. The polymeric oligomers of the invention are derived from these chiral monomers, leading to stereoregular (homochiral) or stereo-alternating polymers (heterochiral). Such stereocontrolled and stereodefined oligomers derived from the chiral monomers of the invention can be used to influence the strand orientations of PNA in PNA: DNA complexes and have benefits in terms of unambiguous, orientation dependent complementary recognition of DNA by the oligomers of the invention.

In a further aspect of the invention, a novel process for the synthesis of the monomers is described (Scheme 1). The route consists of 2 steps starting from 4-hydroxyprolinemethylester hydrochloride and involves N-alkylation, followed by a Mitsunobu reaction. The invention also comprises a process for making the (N-Boc)-2-

aminoethyl bromide, which is the alkylating agent. The process described in the invention is also applicable for the synthesis of the other diastereomers, depending on the stereochemistry of the starting material.

It is observed that oligomers of the present invention overcome some of the biochemical / biophysical limitations of peptide nucleic acids I such as low solubility in water and relative orientations of strands in duplex and triplex structures. The oligomers of the invention are synthesized by a methodology similar to that of PNA, in which the monomer synthons are coupled either to the standard PNA monomer or another modified monomer of the invention. This is useful in adaptation of the present synthetic set-ups for the synthesis of oligomers of the invention.

General method for the synthesis of monomers

One monomer synthon, according to the invention, is prepared by N-alkylation of 4*R*-hydroxy-2*S*-prolinemethyl ester 1 with (N-Boc)-2-aminoethylbromide 2 to afford [1-(N-Boc)-2-aminoethyl]-4*R*-hydroxy-2*S*-proline derivative 3 (Figure 2). The alkylating agent 2 was prepared in two steps from 2-aminoethanol by first making its NH-Boc derivative followed by bromination (Figure 1). The C4 hydroxyl function in 3 was replaced with N3-benzoylthymine using Mitsunobu reaction to yield the (2*S*,4*S*) ester 4a (Figure 3). Similarly, N-alkylation of 4*R*-hydroxy-2*R*-prolinemethylester 5 (Figure 4), followed by Mitsunobu reaction with N3-benzoylthymine (Figure 5) gave the (2*R*,4*S*)-ester 6a. The (2*R*,4*S*) 6a and (2*S*,4*S*) 4a esters were hydrolyzed (Figure 6) to provide the cyclic monomers 4b and 6b for the oligomer assembly.

Additional objects, advantages and novel features of the cyclic monomers, and the process invention will become apparent to those skilled in the art of synthesis for e.g., the 4-hydroxy groups in the starting prolinemethylesters can be inverted by Mitsunobu reaction to obtain the other two diastereomers. Other features of the invention are illustrated by the following examples thereof, which are not intended to be limiting.

General method for synthesis of oligomers of the invention

Oligomers of the invention were prepared generally in accordance with the currently practiced art using solid phase synthesis (Figure 7). Merrifield resin derivatized

with N-Boc- β -alanine (0.13 meq/g of resin) was treated with a three-fold excess of a monomer to be coupled in DMF, along with the coupling agents, diisopropylcarbodiimide and HOBt, (0.4 meq each/g resin) in DMF (2ml/g resin). The removal of the Boc protecting group in the product is accomplished by trifluoroacetic acid treatment and the efficiency of the deprotection and the coupling reactions were determined by the Kaiser test. After sequential addition of monomers, the product oligomer comprising the monomers of the invention was released from the resin by treatment with trifluoroacetic acid-trifluoromethane sulphonic acid. The products were purified using FPLC, their purity checked by HPLC (Figure 9) with acetonitrile-water (0.1% TFA) gradient and their molecular composition confirmed by MALDI-TOF mass spectrometry (Figure 10). The oligomer sequences synthesized by this method are listed in Figure 8.

General method for evaluating the stability of hybrids of compounds under invention with complementary polynucleotides

The oligomers of the invention prepared as above were evaluated for their ability to hybridize with complementary oligonucleotides. This was done by monitoring the temperature dependent UV absorbance (260 nm) of stoichiometrically mixed solutions of PNA oligomers with that of specific sequences of oligodeoxyribonucleotides. The temperatures at the midpoint of the observed sigmoidal transitions (T_m) were taken as a measure of the stability of the PNA: DNA hybrids (Figure 11, Table 1). When the stoichiometry of PNA: DNA is 2:1, the complexes are present as triplexes. The absence of a well-defined transition was considered as evidence for unstable PNA: DNA complexes. The results indicated that the presence of even a single subunit of compounds of the invention II increased the stability of the derived hybrids as compared to the standard PNA of identical sequence (*aeg*PNA I). An increase in the number of monomeric units of the invention further enhanced the stability of complexes to the extent that the homooligomeric PNA: DNA complex did not disassociate even upto 80°C, indicating an extremely high stability. The presence of a single mismatch in the complementary oligonucleotide of PNA: DNA mixture led to very significant destabilization of the complex. The formation of complexes was also checked

independently by gel electrophoresis experiments under non-denaturing conditions (Figure 12).in which the successful complex formation is accompanied by a retardation in the electrophoretic mobility of the corresponding samples. Further, the oligomers or their complexes even with prolonged storage did not precipitate, unlike analogous standard PNA solutions, suggesting remarkably improved aqueous solubility of the oligomers of the invention. These results demonstrate the novel properties of the oligomers composed of the preferred monomers of the invention – namely *enhancing the aqueous solubility and stability of the complexes with oligonucleotides, without compromising the sequence fidelity.*

Those skilled in the art will appreciate that numerous changes and modifications may be made to the preferred embodiments of the invention and that such changes and modifications may be made without departing from the spirit of the invention. It is therefore intended that the appended claims cover all such equivalent variations as fall within the true spirit and scope of the invention.

Table 1: UV- T_m (°C) of DNA:PNA complexes*

	DNA:PNA ₂	2'S	2'R
1	14:7	44	42 (38)
2	14:8	51	54
3	14:9	>75	>75 (67)
4	14:10	>80	>80 (>74)
5	15:9	nd	nd
6	15:10	nd	nd
7	14:11		43 (38)
	DNA:PNA		
8	16:12	41	
9	16:13		39

*nd: not detected. Buffer: 10mM sodium phosphate, pH 7.4. Values in brackets: T_m (°C) with 100 mM NaCl. T_m values are accurate to (\pm) 1°C. Experiments were repeated at least thrice and the T_m values were obtained from the peaks in the first derivative plots.

Another embodiment of this invention provides a method of using peptide nucleic acid oligomers of the present invention for diagnosing and/or modulating the expression of genes in organisms. The modulation includes inhibiting transcription and replication of the said gene.

Yet another embodiment of the present invention provides a process for treating disease conditions associated with undesired protein production in an organism by using the novel compounds of the invention.

Still another embodiment of the present invention provides a pharmaceutical composition comprising a compound of the present invention along with any other pharmaceutically effective agents.

Additional objects, advantages and novel features of this invention will become apparent to those skilled in the art upon examination of the following examples thereof, which are not intended to be limiting.

EXAMPLE 1

Preparation of the precursor (N-Boc)-2-aminoethanol, needed to prepare the alkylating agent.

To a cooled, stirred solution of 2-aminoethanol (5.9ml, 0.098mol) and sodium hydroxide (3.6g, 0.098mol) in water-dioxane (1:1), was added drop-wise Boc-azide (10ml, 0.081mol). The reaction was stirred at room temperature overnight. The dioxane was then removed under vacuum, and the water layer extracted several times with ethyl acetate. The organic layer was dried over sodium sulphate and then, evaporated to dryness under vacuum to get the product (N-Boc)-2-aminoethanol (5.37g, 41% yield), which was used in further steps without purification.

EXAMPLE 2

Preparation of the alkylating agent (N-Boc)-2-aminoethylbromide 2 required for the synthesis of compound 3

To an ice-cooled solution of (N-Boc)-2-aminoethanol (2.5g, 15.5mmol) and carbon tetrabromide (7.5g, 22.7mmol) in dry benzene (25ml), was added triphenyl phosphine (4.7g, 17.9mmol) in small portions over 10-15min. After 30 min. solvent was evaporated and the product immediately purified by silica gel column chromatography to get (N-Boc)-2-aminoethylbromide (5.53g, 79%).

^1H NMR (CDCl_3) δ : 4.95(br s, 1H, NH), 3.50(m, 4H, $(\text{CH}_2)_2$), 1.45(s, 9H, $\text{C}(\text{CH}_3)_3$)

EXAMPLE 3

Alkylation of 4(R)-hydroxy-2(S)-prolinemethylester 1a with 2 to obtain starting material [1-(N-Boc)-2-aminoethyl]-4(R)-hydroxy-2(S)-prolinemethylester 3 for the synthesis of monomers of the invention

A mixture of 4(R)-hydroxy-2(S)-prolinemethylester hydrochloride (3.24g, 17.9mmol), (N-Boc)-2-aminoethylbromide (2.0g, 8.9mmol) and anhydrous potassium carbonate (3.69g, 26.7mmol) was stirred together in DMF: acetonitrile (1:1) at room temperature for 72 h under argon atmosphere. After completion of the reaction as indicated by TLC, solvents were removed *in vacuo*. The residue was taken in water and extracted with ethyl acetate (4x 30ml). The organic layer was dried over sodium sulphate and concentrated to get the crude product, which was purified by silica gel column chromatography. The pure product was obtained in 56% yield.

^1H NMR (CDCl_3) δ : 5.30(br s, 1H, NH), 4.40(m, 1H, H4), 3.70(s, 3H, OCH_3), 3.50(t, 1H, H5), 3.38(dd, 1H, H5'), 3.10(dd, 2H, Boc-NH- CH_2), 2.70(br m, 4H, H2, Boc-NH- CH_2 - CH_2 , OH), 2.50(dd, 1H, H3), 2.10(m, 1H, H3'), 1.40(s, 9H, $\text{C}(\text{CH}_3)_3$)

EXAMPLE 4

Alkylation of 4(R)-hydroxy-2(R)-prolinemethylester 1b with 2 to obtain starting material [1-(N-Boc)-2-aminoethyl]-4(R)-hydroxy-2(R)-prolinemethylester 5 for the synthesis of monomers of the invention

A mixture of 4(R)-hydroxy-2(R)-prolinemethylester hydrochloride (3.24g, 17.9mmol), (N-Boc)-2-aminoethylbromide (2.0g, 8.9mmol) and anhydrous potassium carbonate (3.69g, 26.7mmol) were stirred together in DMF: acetonitrile (1:1) at room temperature

for 72 h under argon atmosphere. After completion of reaction as indicated by TLC, solvents were removed *in vacuo*. The residue was taken in water and extracted with ethyl acetate (4x 30ml). The organic layer was dried over sodium sulphate and concentrated to get the crude product, which was purified by silica gel column chromatography. The pure product was obtained in 56% yield from (N-Boc)-2-aminoethyl bromide.

¹H NMR (CDCl₃) δ: 5.30(br s, 1H, NH), 4.25(br s, 1H, H₄), 3.72(s, 3H, OCH₃), 3.37(m, 1H, H₅), 3.40-3.05(m, 4H, H_{5'}, Boc-NH-CH₂, OH), 2.65(m, 3H, H₂, Boc-NH-CH₂-CH₂), 2.40(m, 1H, H₃), 1.92(dd, 1H, H_{3'}), 1.40(s, 9H, C(CH₃)₃).

¹³C NMR (CDCl₃) δ: 174.9(COOCH₃), 156.1(COOC(CH₃)₃), 78.6(C(CH₃)₃), 70.0(OCH₃), 64.6(C₄), 61.6(C₅), 53.9(C₃), 51.9(C₂), 39.0(Boc-NH-(CH₂)₂), 28.3((CH₃)₃).

EXAMPLE 5

[1-(N-Boc)-2-aminoethyl]-4(S)-(N3-benzoylthymine-1-yl)-2(S)-prolinemethylester 4a

To a stirred solution of [1-(N-Boc)-2-aminoethyl]-4(R)-hydroxy-2(S)-prolinemethylester **3** (0.9g, 3.2mmol), N3-benzoylthymine (0.74g, 3.2mmol) and triphenyl phosphine (1g, 3.8 mmol) in dry THF (10ml) at room temperature, was added dropwise diisopropylazodicarboxylate (DIAD, 0.76ml, 3.8mmol). After completion of the reaction as indicated by TLC (24h), the solvent was removed *in vacuo* and residue purified by silica gel column chromatography to get the pure product (0.6g, 38%).

¹H NMR (CDCl₃) δ: 8.10(s, 1H, T-H₆), 7.90(d, 2H, Bz, *o*-CH), 7.65(t, 1H, Bz-*p*-CH), 7.50(t, 2H, Bz, *m*-CH), 5.25(m, 2H, Boc-NH, H₄), 3.80(s, 3H, OCH₃), 3.25(m, 4H, Boc-NH-CH₂, H₅, H_{5'}), 2.80(m, 3H, Boc-NH-CH₂-CH₂, H₂), 2.60(m, 1H, H₃), 2.00(s, 3H, T-CH₃), 1.95(m, 1H, H_{3'}), 1.45(s, 9H, (CH₃)₃)

EXAMPLE 6

[1-(N-Boc)-2-aminoethyl]-4(S)-(N3-benzoylthymine-1-yl)-2(R)-prolinemethylester 6a

To a stirred solution of 1-(N-Boc-aminoethyl)-4(R)-hydroxy-2(R)-prolinemethylester **5** (0.9g, 3.2mmol), N3-benzoylthymine (0.74g, 3.2mmol) and triphenyl phosphine (1g, 3.8 mmol) in dry THF (10ml) at room temperature, was added dropwise diisopropylazodicarboxylate (DIAD, 0.76ml, 3.8mmol). After completion of

the reaction as indicated by TLC (24h), the solvent was removed *in vacuo* and the residue purified by silica gel column chromatography to get the pure product (0.47g, 30%).

^1H NMR (CDCl_3) δ : 7.92(d, 2H, Bz-*o*-CH), 7.65(t, 1H, Bz-*p*-CH), 7.50(t, 2H, Bz-*m*-CH), 7.40(s, 1H, *T*-H6), 5.25(m, 1H, *H*4), 4.90(br t, 1H, Boc-NH), 3.90(dd, 1H, *H*5), 3.72(s, 3H, OCH_3), 3.30(m, 3H, *H*5', Boc-NH- CH_2), 2.92(dd, 1H, *H*2), 2.78(t, 2H, Boc-NH- CH_2 - CH_2), 2.55(m, 1H, *H*3), 2.20(m, 1H, *H*3'), 2.00(s, 3H, *T*- CH_3), 1.45(s, 9H, $\text{C}(\text{CH}_3)_3$)
 ^{13}C NMR (CDCl_3) δ : 172.5(COOCH_3), 169.0($\text{COOC}(\text{CH}_3)_3$), 162.5(*T*-C2), 155.8(Ph-CO), 149.5(*T*-C4), 137.5(*T*-C6), 134.8, 131.3, 130.0 & 128.9(Ph), 111.1(*T*-C5), 78.8($\text{C}(\text{CH}_3)_3$), 63.1(*C*4), 55.5(Boc-NH- CH_2), 53.7(*C*2), 51.4(OCH_3), 50.6(*C*5), 38.7(Boc-NH- CH_2 - CH_2), 35.1(*C*3), 28.1($\text{C}(\text{CH}_3)_3$), 12.3(*T*- CH_3)

EXAMPLE 7

Solid phase synthesis, deprotection and purification of the oligomers 7 - 13 (Figure 8) of the invention

The oligomers of the invention were synthesized on Merrifield resin (100 mg, pre swollen in dichloromethane) derivatized with β -alanine as a spacer chain (0.13 meq/g of resin). Solid phase synthesis was carried out employing the well known Boc protection strategy involving successive cycles of (i) Boc deprotection using TFA-DCM (1:1, 2 mL, 20 min), (ii) neutralization of the resulting TFA salt by washing with 5% DIPEA in DCM, (2mL, 5 min) (iii) coupling of the desired *aep/aeg*PNA monomer in DMF using HOBT and diisopropylcarbodiimide as the coupling agents (2-5 h). At the end, the oligomers were cleaved from the solid support by TFMSA/TFA to obtain oligomers with β -alanine as a free carboxylic acid at C-terminus. The oligomers 7 - 13 were purified initially by gel filtration over Sephadex G25 and later by reverse phase FPLC on a C-8 column, eluting with an ascending gradient of water-acetonitrile containing 0.1% TFA. The purity was checked by reverse phase HPLC on a C-18 column and molecular composition confirmed by MALDI-TOF mass spectrometric analysis.

EXAMPLE 8

DNA hybridization studies with the oligomers of the invention:

The concentration of oligonucleotides (14 - 16, Figure 8) and oligomers of the invention were determined from the absorbance at 260 nm, using the molar extinction coefficients, of A, $15.4 \text{ cm}^2/\mu\text{mol}$; T $8.8 \text{ cm}^2/\mu\text{mol}$, G, $11.7 \text{ cm}^2/\mu\text{mol}$, C, $7.3 \text{ cm}^2/\mu\text{mol}$.

000260-4479950

All T_m measurements were carried out on Perkin-Elmer Lambda UV spectrometer equipped with a multiple cell holder and variable temperature accessory. A programmable Julabo water circulator controlled the temperature. The samples were prepared by mixing calculated amounts of stock oligonucleotide and the oligomer of the invention together with the addition of calculated amounts of NaCl and sodium phosphate, pH 7.0. to a final volume of 2 mL in a sample cuvette. The samples were annealed by heating at 85°C for 5 min, followed by slow cooling to 5°C. The OD₂₆₀ was then recorded in steps from 10-85°C, with a temperature increment of 0.5°C / min. The results were plotted as percent hyperchromicity at 260 nm as a function of temperature. Analyses of the data were performed by using Microcal Origin software. The melting temperatures were determined from the midpoints of the percent hyperchromicity at 260nm Vs temperature plots.

EXAMPLE 9

Gel electrophoretic experiments

The samples were prepared by mixing stock solutions of the oligomers of the invention and oligonucleotides in tris-EDTA buffer, pH 7.0. These were annealed by heating at 85°C for 5 min., followed by slow cooling to 10°C. The samples were subjected to gel electrophoresis using 15% PAGE (acrylamide: bisacrylamide, 29:1) gels of 1 mm thick and 15 cm long. Electrophoresis were carried out at a constant voltage of 150 V till the marker bromophenol dye migrated to 12 cm length on the gel. The bands were visualized by illuminating the gel (254 nm) after placing it on a fluorescent thin-layer silica gel chromatographic plate.

SEQUENCE LISTING

(1) GENERAL INFORMATION

- (i) APPLICANT:
- (ii) TITLE OF INVENTION
- (iii) NUMBER OF SEQUENCES
- (iv) CORRESPONDENCE ADDRESS
- (v) COMPUTER READABLE FORM
 - (A) MEDIUM TYPE: Floppy Disk
 - (B) COMPUTER: IBM PC Compatible
 - (C) OPERATING SYSTEM: MS-DOS
 - (D) SOFTWARE: Microsoft Word
- (vi) CURRENT APPLICATION DATA
- (vii) ATTORNEY / AGENT INFORMATION
- (viii) TELECOMMUNICATION INFORMATION

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS
 - (A) LENGTH: 8
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: Unknown
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 1
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 2
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 3
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 4
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 - (A) NAME / KEY : Modified site

/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 4
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 5
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 6
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 7
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 8
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO :2:
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa β -Ala
1 5 8

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS
(A) LENGTH:
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: Unknown
- (iv) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 1
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:

- (A) NAME / KEY : Modified site
 (B) LOCATION: 2
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 3
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 4
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 5
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
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 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 7
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 8
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO :3:
- | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-------|
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| 1 | | | | | 5 | | | 8 | |
- (2) INFORMATION FOR SEQ ID NO:4:
- (i) SEQUENCE CHARACTERISTICS
 (A) LENGTH: 8
 (B) TYPE: amino acid

- (C) STRANDEDNESS: single
(D) TOPOLOGY: Unknown
- (v) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 1
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 2
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 3
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 4
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 5
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 6
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 7
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 8
(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO :4:

Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	β-Ala
1					5			8	

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS

(A) LENGTH: 8

(B) TYPE: amino acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: Unknown

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 1

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 2

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 3

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 4

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 5

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 6

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 7
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 8
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO :2:

Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	β-Ala
1					5			8	

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: Unknown
- (vi) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 1
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 2
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 3
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."

- (ix) FEATURE:
 - (A) NAME / KEY : Modified site
 - (B) LOCATION: 4
 - (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
 - (A) NAME / KEY : Modified site

- (B) LOCATION: 5
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 7
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
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- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 8
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 9
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 10
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 11
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 12
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO :6:
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa-β-Ala
 1 5 10

(2) INFORMATION FOR SEQ ID NO:7:

(i) SEQUENCE CHARACTERISTICS

- (A) LENGTH: 12
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: Unknown

(vii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME / KEY : Modified site
- (B) LOCATION: 1
- (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
- (B) LOCATION: 2
- (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
- (B) LOCATION: 3
- (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
- (B) LOCATION: 4
- (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
- (B) LOCATION: 5
- (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
- (B) LOCATION: 6
- (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
- (B) LOCATION: 7
- (D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
 (B) LOCATION: 8
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 9
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 10
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 11
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 12
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO :6:
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa-β-Ala
 1 5 10
- (2) INFORMATION FOR SEQ ID NO:8:
 (i) SEQUENCE CHARACTERISTICS
 (A) LENGTH: 12
 (B) TYPE: amino acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: Unknown
 (ii) MOLECULE TYPE: peptide
 (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 1
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 2

- (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N9 of the adenine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 3
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
 (A) NAME / KEY : Modified site
 (B) LOCATION: 4
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N9 of the adenine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 5
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 7
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 8
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N9 of the adenine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 9
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 10
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N9 of the adenine."

- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 11
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."

- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 12
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = " Thymine heterocycle is attached to N1-(2-aminoethyl) proline at C4 of proline through position 1 of the heterocycle."

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO :9:
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa β-ala
 1 5 10

(2) INFORMATION FOR SEQ ID NO: 10:

(i) SEQUENCE CHARACTERISTICS

- (A) LENGTH: 12
 (B) TYPE: amino acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: Unknown

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME / KEY : Modified site
 (B) LOCATION: 1
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
 (B) LOCATION: 2
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
 (B) LOCATION: 3
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = " Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(A) NAME / KEY : Modified site

(B) LOCATION: 4

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Cytosine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the cytosine."

(ix) FEATURE:

- (A) NAME / KEY : Modified site
 (B) LOCATION: 5
 (D) OTHER INFORMATION: /label = MODIFIED-SITE

note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 6
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Cytosine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the cytosine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 7
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 8
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Adenine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N9 of the adenine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 9
(D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 10
(D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 11
(D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 12
(D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 13
(D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 14
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 15
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO :10:
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 Xaa β-ala
 1 5 10

INFORMATION FOR SEQ ID NO: 11:

- (i) SEQUENCE CHARACTERISTICS
 (A) LENGTH: 12
 (B) TYPE: amino acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: Unknown
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 1
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 2
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 3
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (A) NAME / KEY : Modified site
 (B) LOCATION: 4
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Guanine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N9 of the guanine."
- (ix) FEATURE:

- (A) NAME / KEY : Modified site
 (B) LOCATION: 5
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Guanine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the guanine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 7
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 8
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N9 of the adenine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 9
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 10
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 11
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 12
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 note = "Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 13

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- (D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 14
(D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 15
(D) OTHER INFORMATION: /label = MODIFIED-SITE
note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO :10:
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
Xaa β -ala
1 5 10

INFORMATION FOR SEQ ID NO:12:

- (i) SEQUENCE CHARACTERISTICS
(A) LENGTH: 10
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: Unknown
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 1
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Guanine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 2
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 3
(D) OTHER INFORMATION: /label = MODIFIED-SITE
/note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
(A) NAME / KEY : Modified site
(B) LOCATION: 4

- (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Guanine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 5
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 7
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 8
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 7 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 9
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."
- (ix) FEATURE:
 (A) NAME / KEY : Modified site
 (B) LOCATION: 10
 (D) OTHER INFORMATION: /label = MODIFIED-SITE
 /note = "Thymine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the thymine."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO :12:
- | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-------|
| Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | Xaa | β-Ala |
| 1 | | | | | 5 | | | | 8 | | |
- INFORMATION FOR SEQ ID NO:13:
- (i) SEQUENCE CHARACTERISTICS
 (A) LENGTH: 10
 (B) TYPE: amino acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: Unknown

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 1

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Guanine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 2

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 3

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 4

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Guanine heterocycle is attached at C4 of N1-(2-aminoethyl) proline through position N1 of the guanine."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 5

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 9 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 6

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 7

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 8

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = "Adenine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 7 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 9

(D) OTHER INFORMATION: /label = MODIFIED-SITE

/note = " Cytosine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(ix) FEATURE:

(A) NAME / KEY : Modified site

(B) LOCATION: 10

(D) OTHER INFORMATION: /label = MODIFIED-SITE

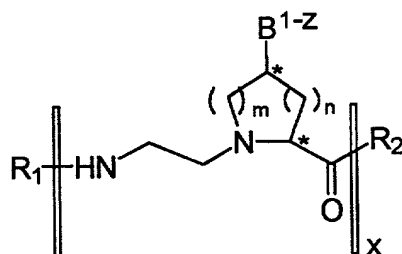
/note = "Thymine heterocycle is attached to N-acetyl (2-aminoethyl) glycine through the N-acetyl group at position 1 of the heterocycle."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO :2:

Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	β-Ala
1					5				8			

CLAIMS:

1. A novel chiral, peptide nucleic acid oligomers having the formula :

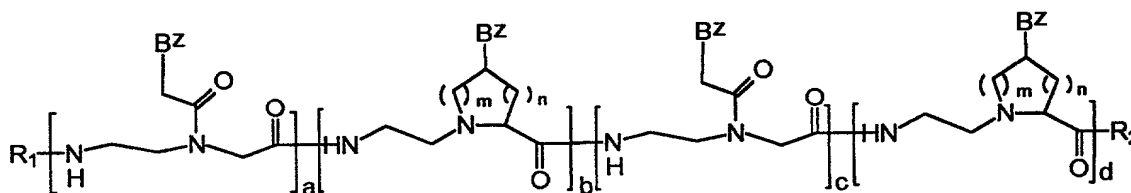


aep PNA II

Wherein,

- m and n are 1 to 2 and x = 1-20;
- each of B¹-B^z is independently selected from the group consisting of H, HO, NH₂, naturally occurring nucleobases adenine (A), thymine (T), cytosine (C) and guanine (G), non-naturally occurring nucleobases, DNA intercalators, heterocyclic moieties and reporter ligands;
- each chiral monomeric unit independently selected from the four possible diastereomers; and
- R₁=H/Fluorophore/Biotin; R₂=OH/NH(CH₂)₂COOH/
NH(CH₂)₃NH(CH₂)₄NH(CH₂)₃NH₂.

2. A novel chiral, peptide nucleic acid oligomers having the formula :



aep PNA III

which are heteropolymeric aepPNA III (with all four possible diastereomers) involving one or more substitution of the non-chiral aeg unit of aminoethylglycyl PNA I in aepPNA II as below:

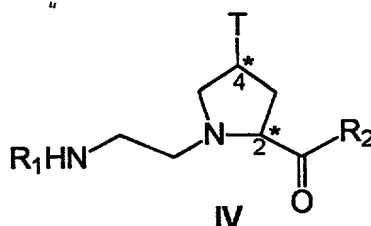
- each chiral monomer unit independently selected from the four possible diastereomers,
- a,b,c,d,m,n are integers with independent values in the range 1 to 10 and various combinations thereof,
- R_1 is $H/COCH_3$ or L (corresponding to a fluorophore e.g. dansyl, carboxyfluorescein),
- R_2 is OH, NH_2 , $NHCH_2CH_2COOH$, sperminyl i.e., $NH(CH_2)_3NH(CH_2)_4NH(CH_2)_3NH_2$, and
- each of B^1-B^z is independently selected from the group consisting of H, HO, NH_2 , naturally occurring nucleobases, non-naturally occurring nucleobases, DNA intercalators, heterocyclic moieties and reporter ligands.

3. A novel chiral, peptide nucleic acid oligomers as claimed in claim 2, wherein $m=n=1$; $B^z = T$; $R_1=H$; $R_2 = NH(CH_2)_2COOH$, with

- i. $a=7, b=1, c=d=0$,
- ii. $a=c=3, b=d=1$,
- iii. $a=b=c=d=1$, repeating twice in that order,
- iv. $a=b=c=0, d=8$,
- v. $a=d=0, b=1, c=7-11$ and with various combinations of B^z .

4. A novel chiral, peptide nucleic acid as claimed in claim 1 and 2, wherein the oligomers are synthesized by adaptation of standard peptide synthesis procedures, either in solution or in solid phase.

5. A monomer precursor-synthon having the formula IV



Wherein,

- $R_1 = H/Boc/Fmoc$, $R_2 = OMe/OH/OEt/Obenzyl$,
 - variation of chirality at positions 2 and 4 leading to four diastereomers (*2S,4R*), (*2R,4S*), (*2S,4S*) and (*2R,4R*), and
 - T is the nucleobase.
6. A monomer precursor-synthon as claimed in claim 5, wherein T is a naturally occurring nucleobase.
7. A process for preparing compounds according to claim 5, comprising the steps of :
- A. providing the alkylating reagent (N-Boc)-2-aminoethylbromide (2) in two steps from 2-aminoethanol;
 - B. providing N-alkylation of 4-hydroxyprolinemethylester with reagent prepared as in claim 7A
 - alkylation of 4*R*-hydroxy-2*S*-prolinemethylester (1a) with (N-Boc)-2-aminoethyl bromide (2) to afford [1-(N-Boc)-2-aminoethyl]-4*R*-hydroxy-2*S*-prolinemethyl ester (3),
 - alkylation of 4*R*-hydroxy-2*R*-prolinemethylester (1b) with (N-Boc)-2-aminoethyl bromide (2) to afford [1-(N-Boc)-2-aminoethyl]-4*R*-hydroxy-2*R*-prolinemethyl ester (5),
 - alkylation of 4*S*-hydroxy-2*R*-prolinemethylester with (N-Boc)-2-aminoethyl bromide (2) to afford [1-(N-Boc)-2-aminoethyl]-4*S*-hydroxy-2*R*-proline methylester
 - alkylation of 4*S*-hydroxy-2*S*-prolinemethylester with (N-Boc)-2-aminoethyl bromide (2) to afford [1-(N-Boc)-2-aminoethyl]-4*S*-hydroxy-2*S*-proline methylester,
 - C. producing monomer synthons (4a) and (6a) by Mitsunobu reaction of compounds prepared according to claim 7B with N3-benzoylthymine.

- 000260-44-092000
8. A process for introducing novel chiral monomers as claimed in claim 7C at specific/desired position(s) in the oligomers of desired sequences
 9. A process for sequence specific recognition of a single or double stranded polynucleotide (DNA, RNA), by the oligomers as per claims 1 and 2 derived from the compounds according to claim 7.
 10. A method of using peptide nucleic acid oligomers as claimed in claim 9 for diagnosing and/or modulating the expression of genes in organisms.
 11. A method as claimed claim 10 wherein said modulation includes inhibiting transcription and replication of the said gene.
 12. A process for treating disease conditions associated with undesired protein production in an organism by using the compound according to claims 1 and 2.
 13. A pharmaceutical composition comprising a compound according to claims 1 and 2 along with any other pharmaceutically effective agents.

ABSTRACT

A novel class of peptide nucleic acid compounds, which are positively charged, chiral analogs based on cyclic monomeric units, bind to complementary ssDNA, dsDNA and RNA strands with more affinity than the corresponding achiral, linear peptide nucleic acids, without compromising the binding sequence specificity and the compounds of this invention are more soluble in water than the standard peptide nucleic acids.

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Figure 1

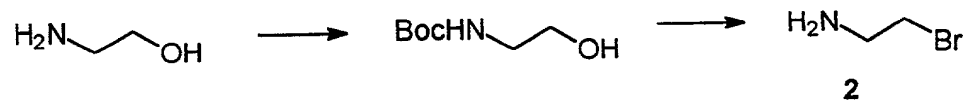


Figure 2

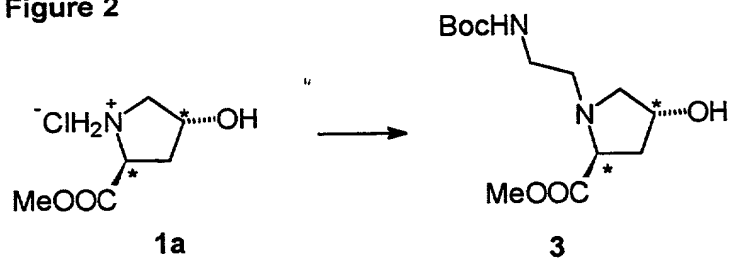


Figure 3

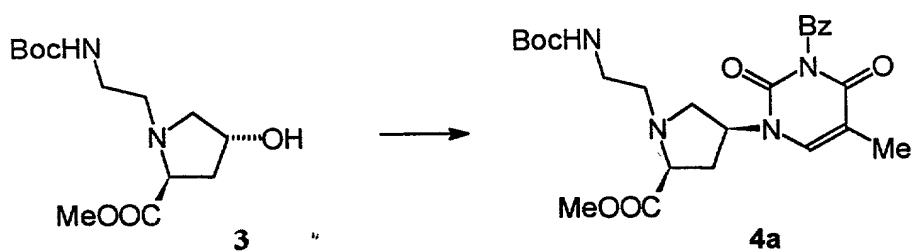


Figure 4

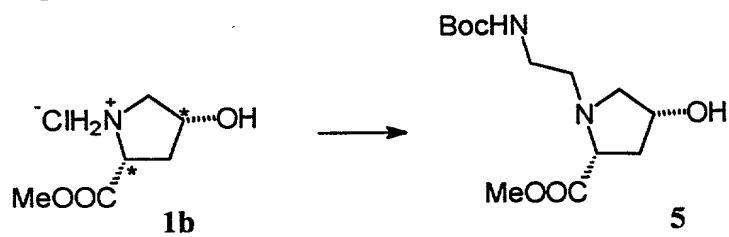


Figure 5

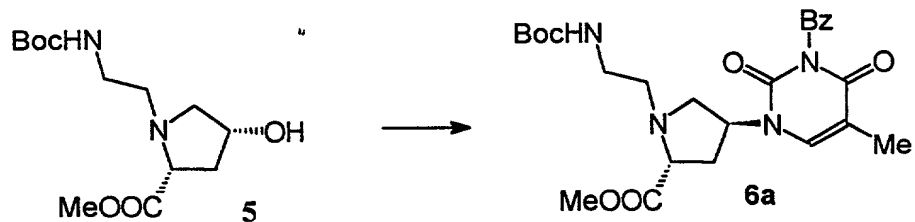


Figure 6

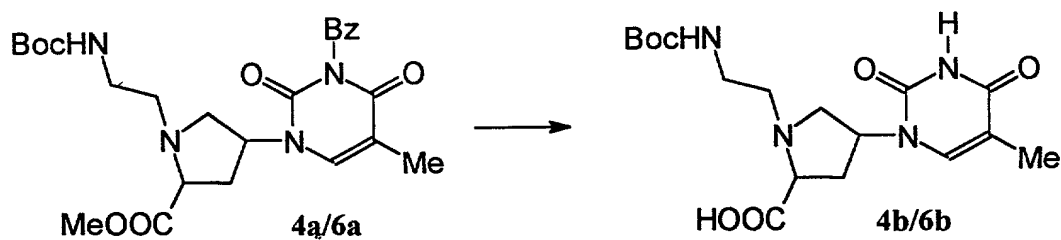


FIG. 6

Figure 7 Solid Phase synthesis of Oligomers

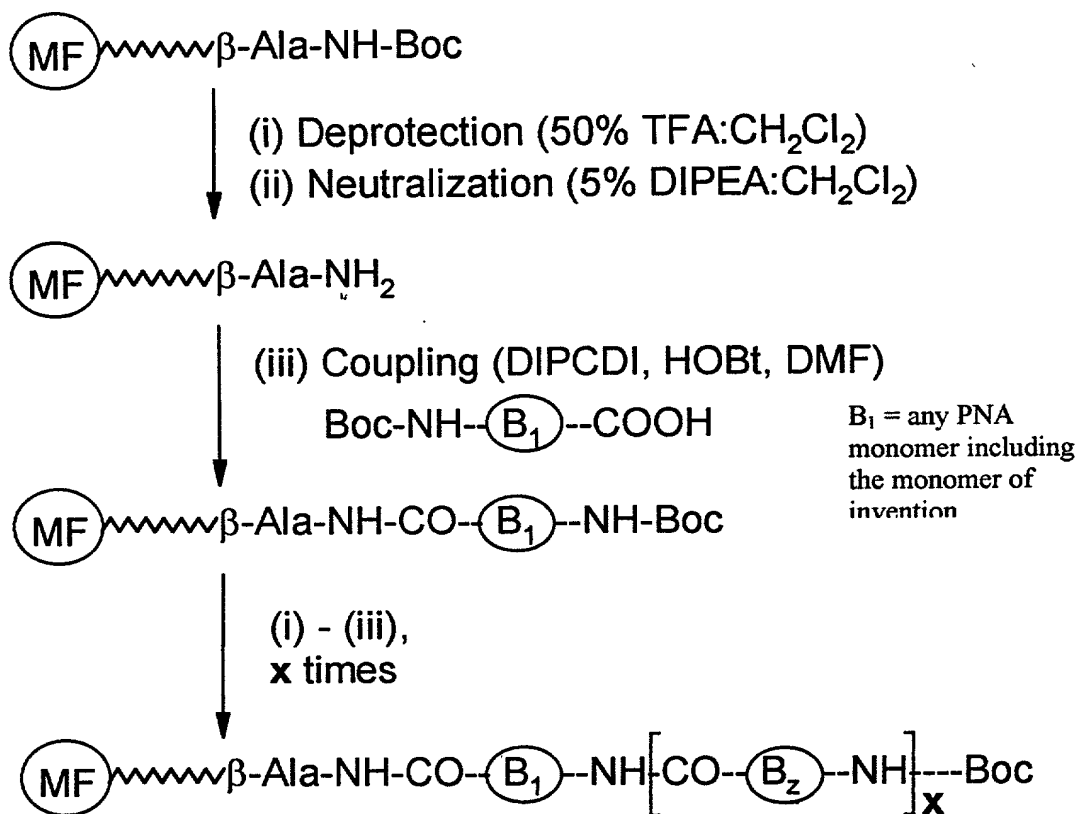


FIG. 7

Figure 8. Oligomer sequences comprising novel monomers of the invention

Oligomer sequences

- 7 H- T T T T T T T t $-(\beta\text{-Ala})\text{-OH}$
 8 H- T T T t T T T t $-(\beta\text{-Ala})\text{-OH}$
 9 H- T t T t T t T t $-(\beta\text{-Ala})\text{-OH}$
 10 H- t t t t t t t t $-(\beta\text{-Ala})\text{-OH}$
 11 H- T T T T T T T T $-(\beta\text{-Ala})\text{-OH}$
 12 H- t A T A T T A T T A T T $-(\beta\text{-Ala})\text{-OH}$
 13 H- T A T A T T A T T A T T $-(\beta\text{-Ala})\text{-OH}$

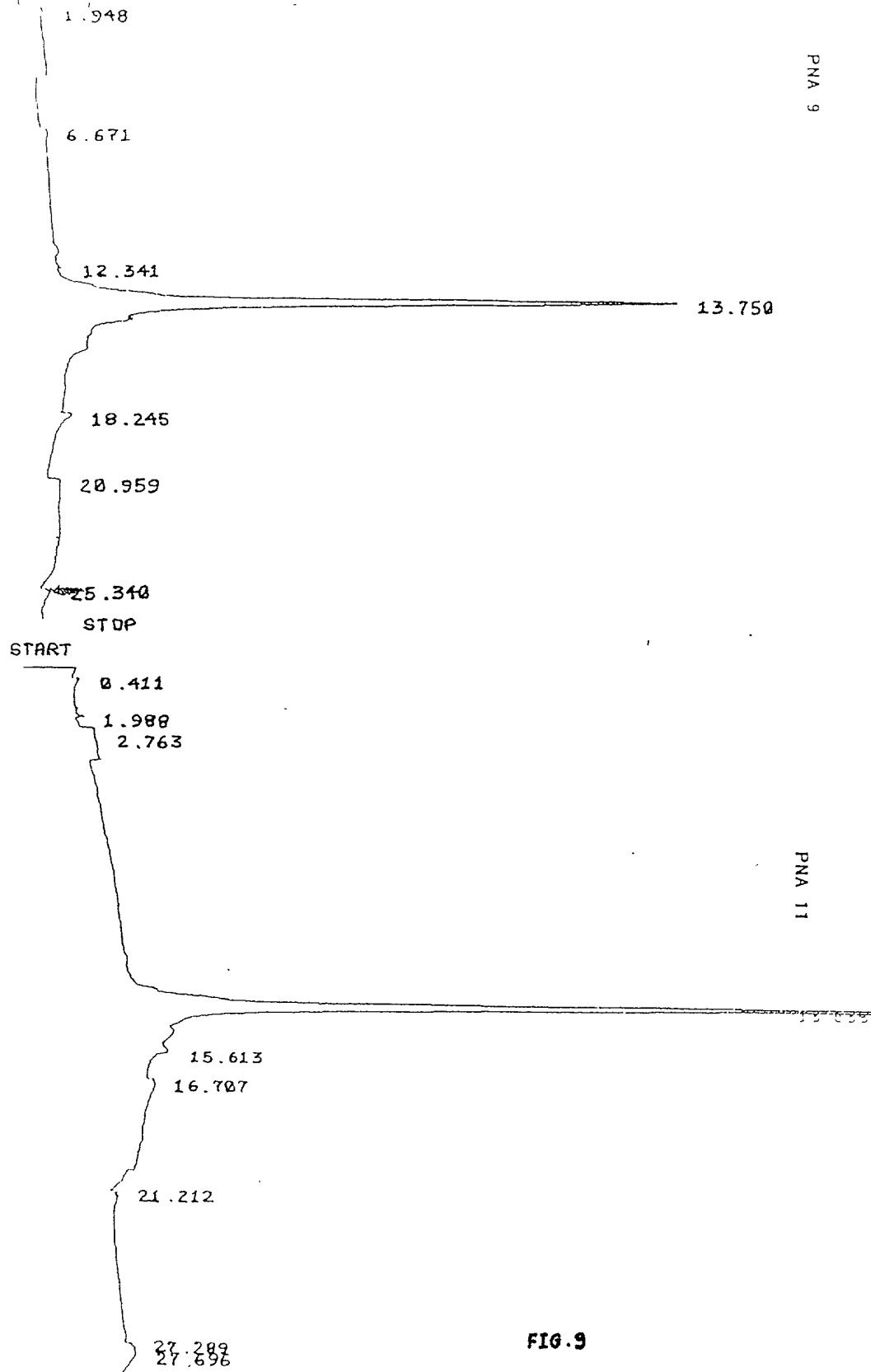
A/T = *aeg*PNA-A/T, t = *aep*PNA -T

DNA sequences

- 14 5'- G C A A A A A A A C G -3'
 15 5'- G C A A A T A A A C G -3'
 16 5'- A A T A A T A A T A T A -3'

FIG. 8

Figure 9a



0002044-092000

Figure 9a

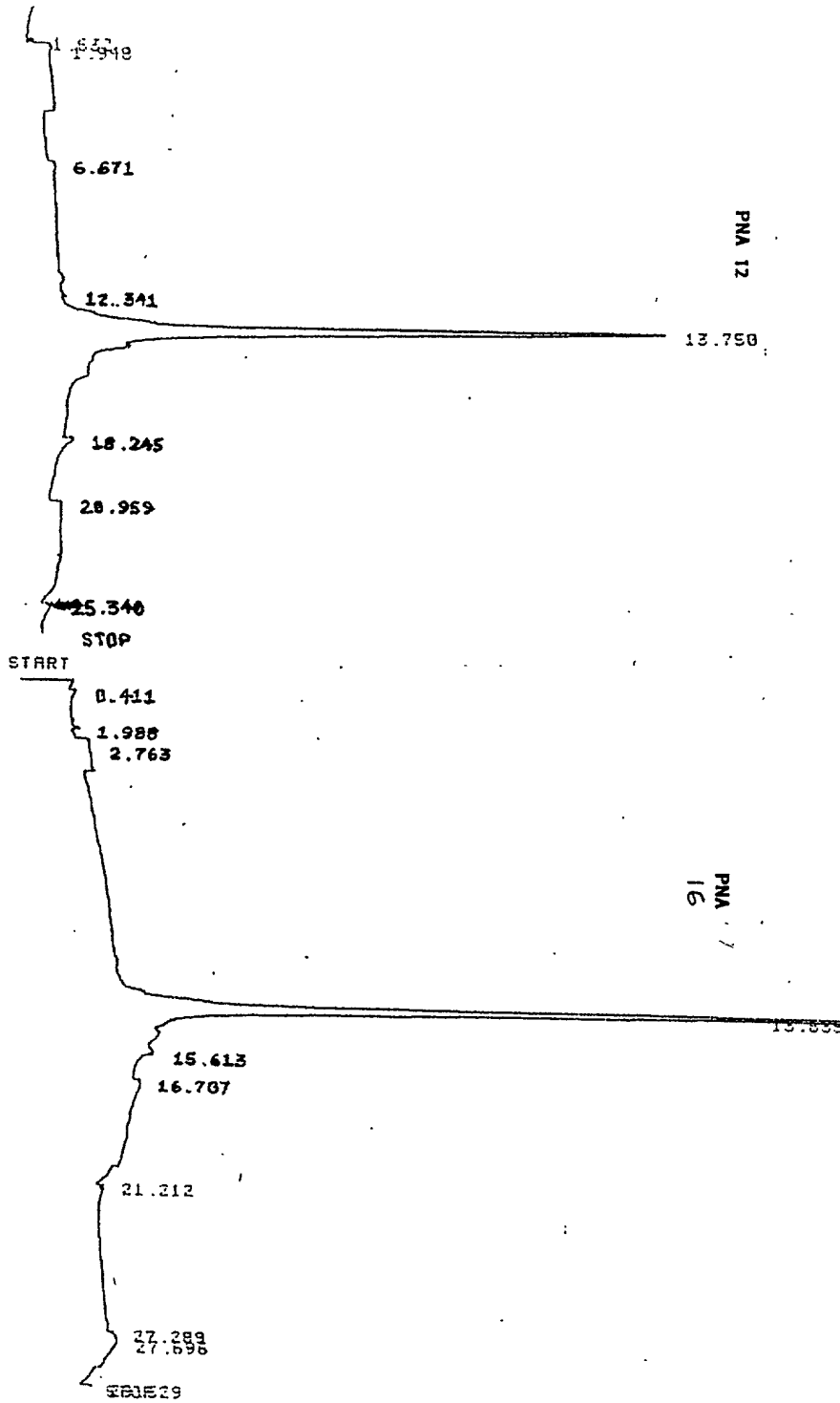


FIG. 9

0905144.092000

Figure 9b

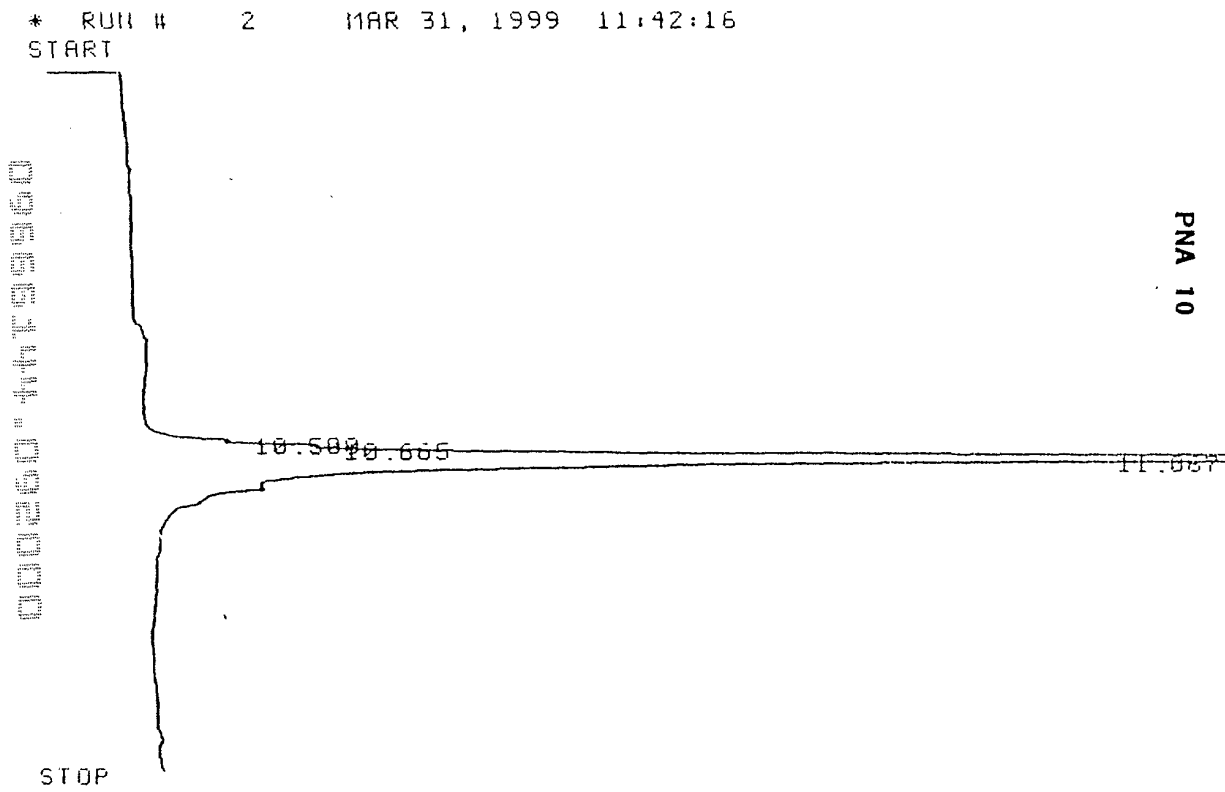


FIG.9

Closing signal file M:SIGNAL .BNC

* RUN # 2 MAR 31, 1999 11:42:16
START

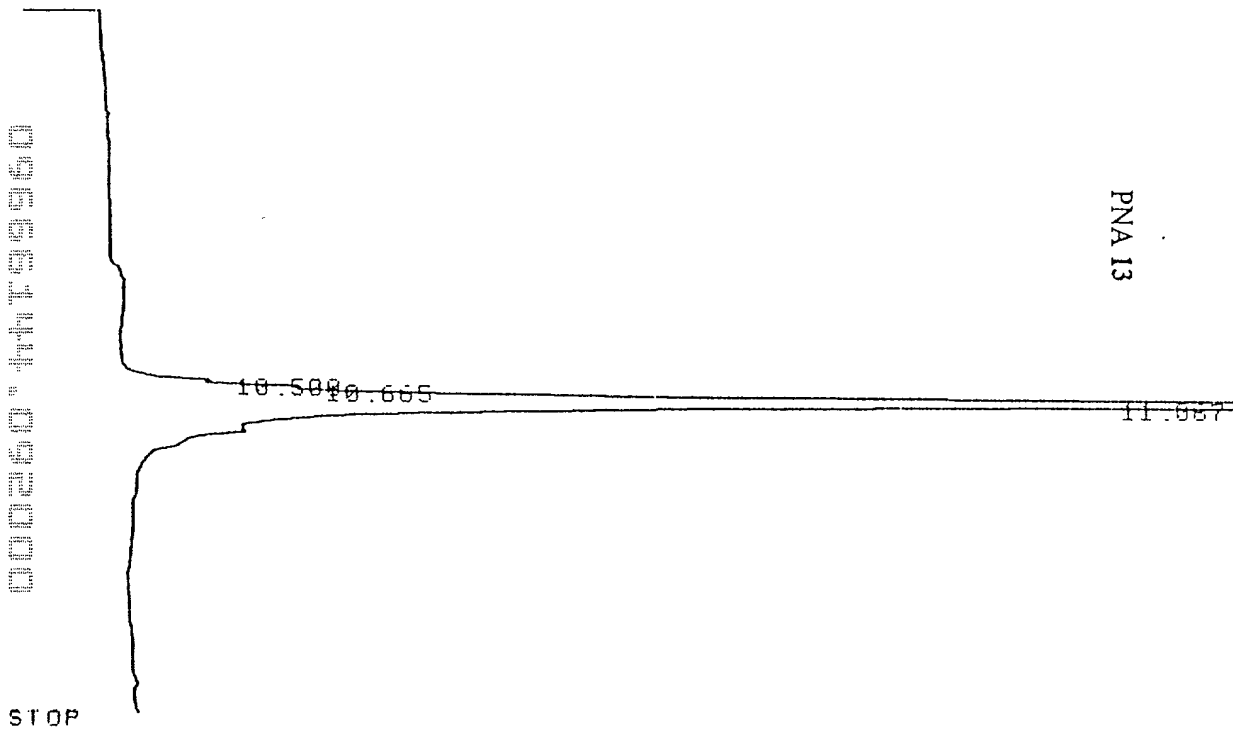


FIG.9

Closing signal file H:SIGNAL .BNC

FIGURE 10 A. MALDI-TOF mass spectrum of PNA 7

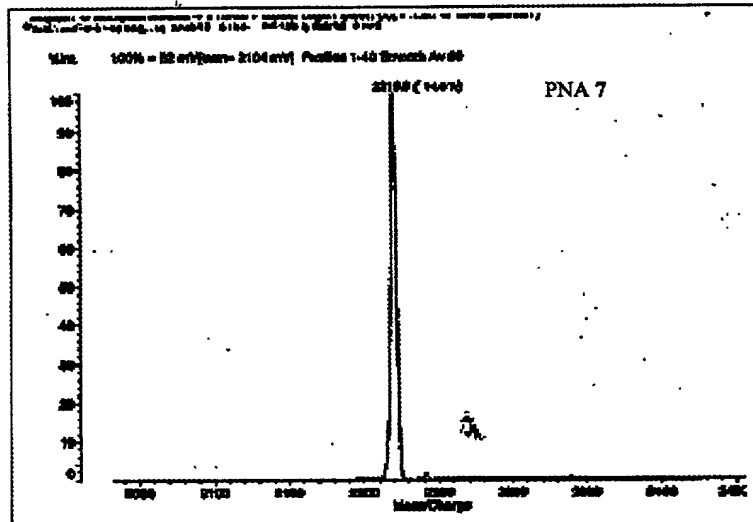


FIG.10

1



FIG. 10

FIGURE 10 A. MALDI-TOF mass spectrum of PNA 10

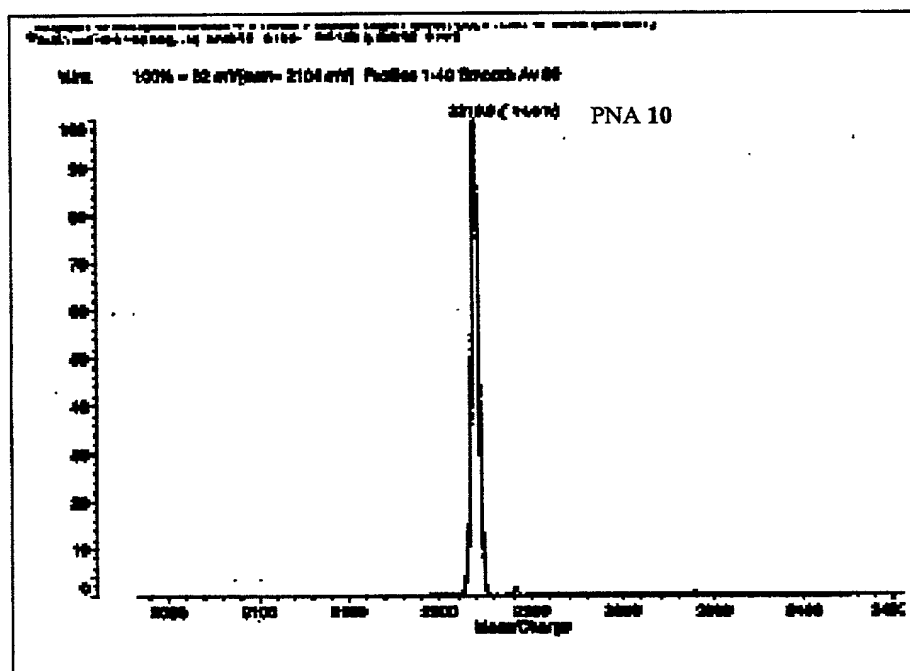


FIG.10

Figure 10B. MALDI-TOF mass spectrum of PNA 11, 12

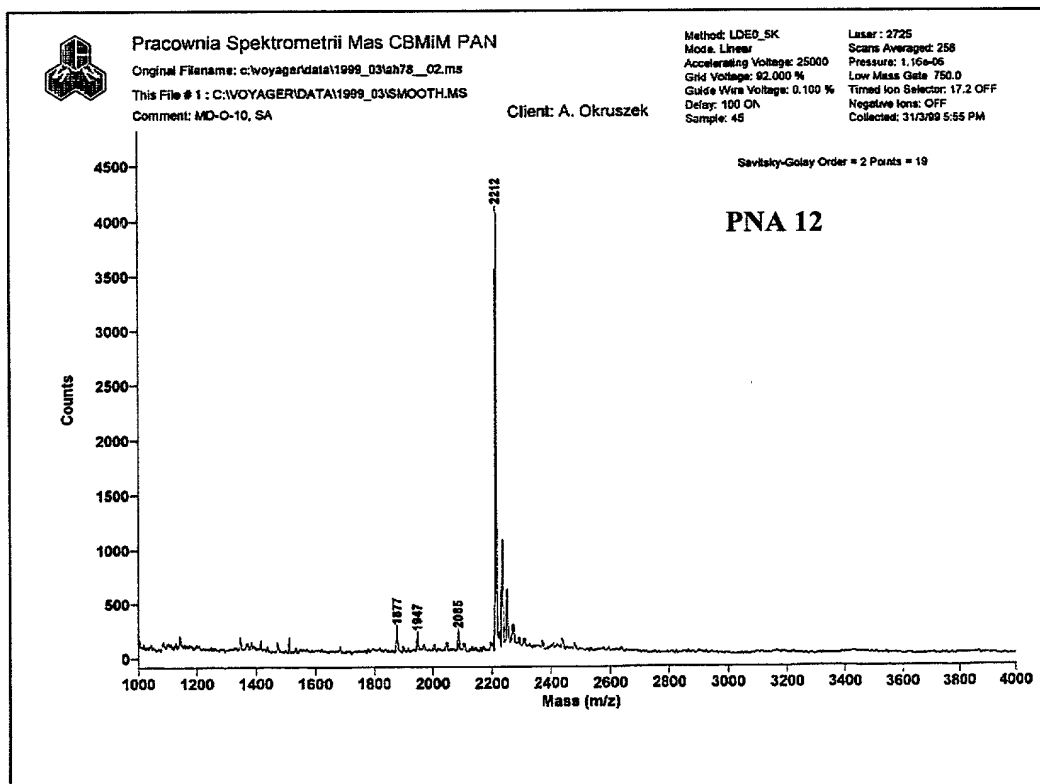
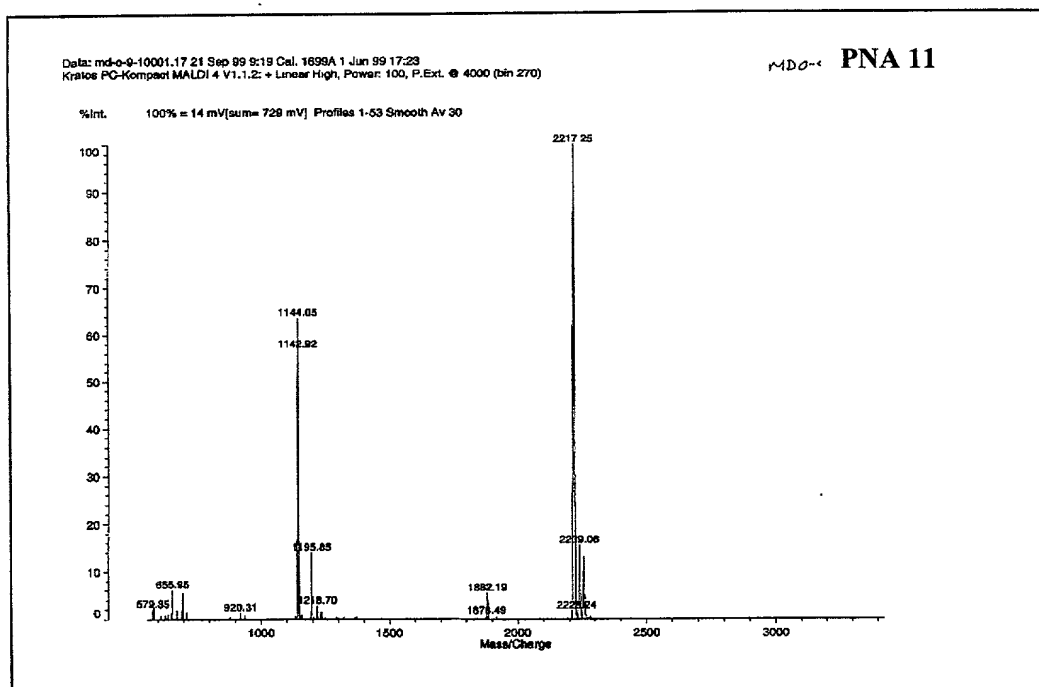
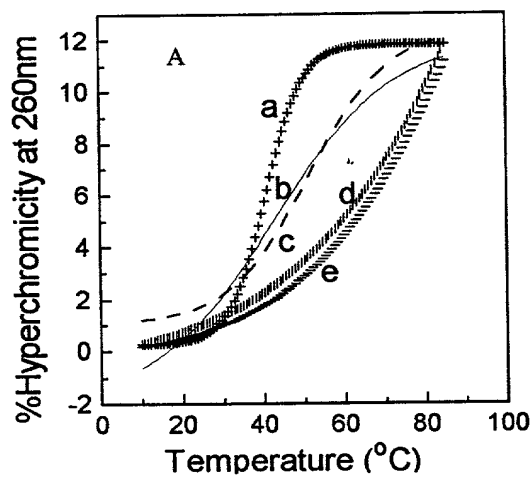


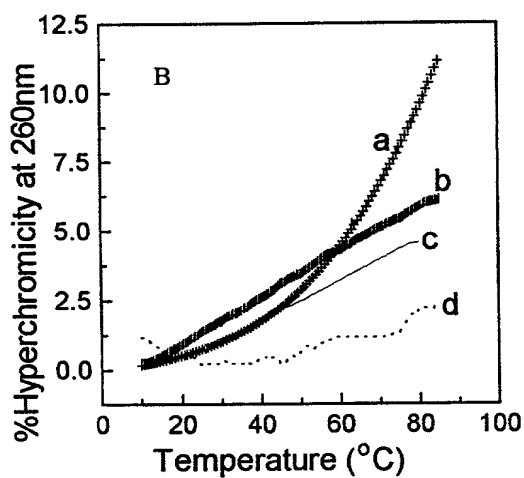
FIG. 10

Figure 11

UV melting profiles



A: for the 2S modification a: 14:11, b: 14:7, c: 14:8, d: 14:9, e: 14:10.

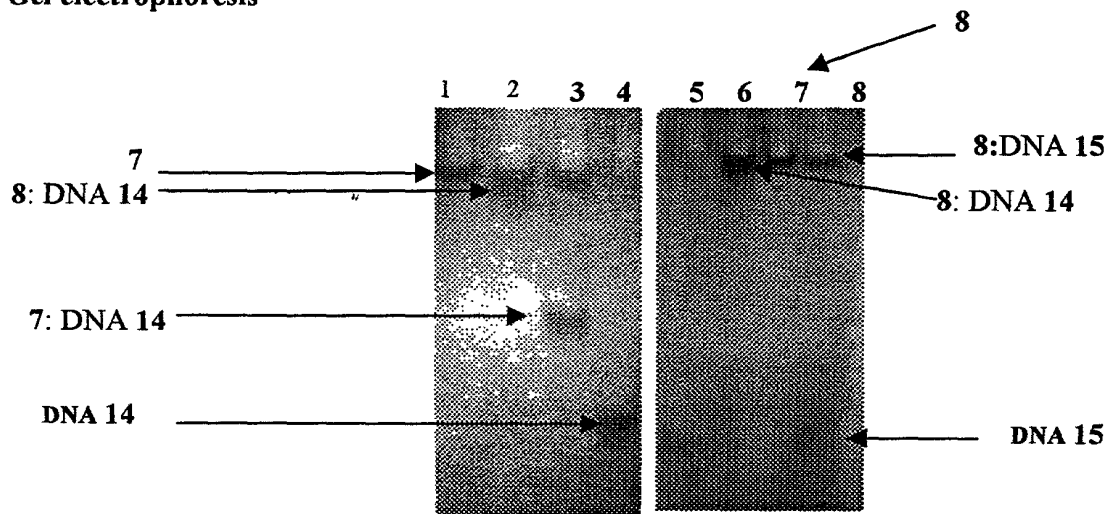


B: a: 14:10, b: 15:10 and of single strands c: 15, d: 10.

FIG.11

FIGURE 12

Gel electrophoresis



15% Polyacrylamide Gel Electrophoresis (acrylamide:bis-acrylamide, 29:1) of *aep*PNA:DNA complexes. Bands were visualized by UV-shadowing, i.e., by illuminating the gel placed on a fluorescent thin-layer silica gel chromatographic plate, F₂₅₄, 20cm x 20cm using UV light. Lane 1: 7; Lane 2: (8: DNA 14); Lane 3: (7: DNA 14); Lane 4: DNA 14; Lane 5: DNA 15; Lane 6: (8: DNA 14); Lane 7: 8; Lane 8: (8: DNA 15).

Oligomer sequences

7 H- T T T T T T T t -(β-Ala)-O-

8 H- T T T t T T T t -(β-Ala)-O-

T = *aep*PNA-T, t = *aep*PNA -T

DNA sequences

14 5'- G C A A A A A A A C G -3'

15 5'- G C A A A T A A A A C G -3'

FIG.12